Sequence of the pYfpGGfp fusion protein.

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_	GTTTGACAGC	TTATCATCGA	CTGCACGGTG	CACCAATGCT	reredectea	GGCAGCCATC	09
61	GGAAGCTGTG	GTATGGCTGT	GCAGGTCGTA	AATCACTGCA	TAATTCGTGT	CGCTCAAGGC	120
121	GCACTCCCGT	TCTGGATAAT	GTTTTTGCG	CCGACATCAT	AACGGTTCTG	GCAAATATTC	180
181	TGAAATGAGC	TGTTGACAAT	TAATCATCCG	GCTCGTATAA	TGTGTGGAAT	TGTGAGCGGA	240
241	TAACAATTTC	ACACAGGAAA	CAGCGCCGCT	GAGAAAAAGC	GAAGCGGCAC	TGCTCTTTAA	300
301	CAATTTATCA	GACAATCTGT	GIGGGCACIC	GACCGGAATT	ATCGATTAAC	TTTATTAT	360
361	AAAATTAAAG	AGGTATATAT	TAATGTATCG	ATTAAATAAG	GAGGAATAAA	CCATGGtgag	420
421	caagggcgag	gagetgttea	coggggtggt	gcccatcctg	gtegagetgg	acggcgacgt	480
481	aaacggccac	aagttcagcg	tgtccggcga	gggcgagggc	gatgccacct	acggcaaget	540
541	gaccctgaag	ttcatctgca	ccaccggcaa	getgeeegtg	ccctggccca	cectegtgae	009
601	caccttegge	tacggcctgc	agtgettege	cagataacaa	gaccacatga	agcagcacga	099
661	cttcttcaag	teegecatge	ccgaaggeta	cgtccaggag	cgcaccatct	tetteaagga	720
721	cgacggcaac	tacaagaccc	gegeegaggt	gaagttcgag	ggcgacaccc	tggtgaaccg	780
781	categagetg	aagggcatcg	acttcaagga	ggacggcaac	atcctggggc	acaagorgga	840
841	gtacaactac	aacagccaca	acgictatat	catggccgac	aagcagaaga	acggcatcaa	006
901	ggtgaacttc	aagatoogoo	acaacatoga	ggacggcagc	gtgcagctcg	cegaceacta	096
961	ccagcagaac	acccccatcg	gcgacggccc	cgtgctgctg	cccgacaacc	actacctgag	1020
1021	ctaccagtcc	gccctgagca	аадассссаа	cgagaagcgc	gatcacatgg	toctgotgga	1080
1081	gttcgtgacc	gccgccggga	tcactctcgg	catggacgag	ctgtacaagA	cragr getga	1140

1141	tactegeatt	ggtgtaacaa	tctataagta	cgacgataac	tttatgtotg	tagtgegeaa	1200
1201	ggctattgag	caagatgega	aagccgcgcc	agatgttcag	ctgctgatga	atgattctca	1260
1261	gaatgaccag	tccaagcaga	acgatcagat	cgacgtattg	ctggccaagg	gggtgaaggc	1320
1321	actggccatc	aacctggttg	acccggcagc	tgegggtaeg	gtgattgaga	aagcgcgtgg	1380
1381	gcaaaacgtg	ccggtggtt	tottcaacaa	agaaccgtct	cgtaaggege	tggatageta	1440
1441	cgacaaagcc	tactacgttg	gcactgactc	aaaagagtee	ggcattattc	aaggcgattt	1500
1501	gattgctaaa	cactgggcgg	cgaatcaggg	ttgggatetg	aacaaagacg	gtcagattca	1560
1561	gttcgtactg	ctgaaaggtg	aaccgggcca	tooggatgea	gaagcacgta	ccacttacgt	1620
1621	gattaaagaa	ttgaacgata	aaggcatcaa	aactgaacag	ttacagttag	ataccgcaat	1680
1681	gtgggacacc	geteaggega	aagataagat	ggacgcctgg	ctgtctggcc	cgaacgccaa	174C
1741	caaaatcgaa	gtggttatcg	ccaacaacga	tgegatggea	atgggcgcgg	ttgaagcgct	1800
1801	gaaagcacac	aacaagteca	gcattccggt	gtttggcgtc	gatgegetge	cagaagcgct	1860
1861	ggagatagta	aaatccggtg	cactggcggg	caccgtactg	aacgatgota	acaaccaggc	1920
1921	gaaagcgacc	tttgatotgg	cgaaaaacct	ggeegatggt	aaaggtgcgg	ctgatggcac	1980
1981	caactggaaa	ategacaaca	aagtggtccg	cgtaccttat	gttggcgtag	ataaagacaa	204C
2041	cctggctgaa	ttcagcaaga	aaGGTACCag	taaaggagaa	gaacttttca	ctggagttgt	2100
2101	cccaattett	gttgaattag	atggtgatgt	taatgggcac	asattttetg	tcagtggaga	2160
2161	gggtgaaggt	gatgcaacat	acggaaaact	tacccttaaa	tttatttgca	ctactggaaa	2220
2221	actacctgtt	ccatggccaa	cacttgtcac	tactttctct	tatggtgttc	aatgettte	2280
2281	ccgttatccg	gatcatatga	aacggcatga	ctttttcaag	agtgccatgc	ccgaaggtta	2340

2341	tgtacaggaa	cgcactatat	ctttcaaaga	tgacgggaac	tacaagacgc	gtgctgaagt	2400
2401	caagtttgaa	ggtgataccc	ttgttaatcg	tatogagtta	aaaggtattg	attttaaaga	2460
2461	agatggaaac	atteteggae	acaaactega	gtacaactat	aactcacaca	atgtatacat	2520
2521	cacggcagac	aaacaaaaga	atggaatcaa	agetaaette	aaattegee	acaacattga	2580
2581	agatggatcc	gttcaactag	cagaccatta	tcaacaaaat	actccaattg	gcgatggccc	2640
2641	tgtcctttta	ccagacaacc	attacctgtc	gacacaatct	gccctttcga	aagatcccaa	270 C
2701	cgaaaagcgt	gaccacatgg	tecttettga	gtttgtaact	getgetggga	ttacacatgg	2760
2761	catggatgag	ctctacaaat	aa aagci tac	GTAGAACAAA	AACTCATCTC	AGAAGAGGAT	2820
2821	CTGAATAGCG	CCGTCGACCA	TCATCATCAT	CATCATIGAG	TTTAAACGGT	CTCCAGCTTG	2880
2881	GCTGTTTGG	CGGATGAGAG	AAGATTTTCA	GCCTGATACA	GATTAAATCA	GAACGCAGAA	294C
2941	GCGGTCTGAT	AAAACAGAAT	TTGCCTGGCG	GCAGTAGCGC	GGTGGTCCCA	CCTGACCCCA	3000
3001	TGCCGAACTC	AGAAGTGAAA	CCCCCTAGCG	CCGATGGTAG	reredeerer	CCCCATGCGA	3060
3061	GAGTAGGGAA	CTGCCAGGCA	TCAAATAAAA	CGAAAGGCTC	AGTCGAAAGA	CIGGGCCTIT	3120
3121	CGTTTTATCT	GTTGTTTGTC	GGTGAACGCT	CTCCTGAGTA	GGACAAATCC	GCCGGGAGCG	3180
3181	GATTTGAACG	TTGCGAAGCA	ACGGCCCGGA	GGGTGGCGGG	CAGGACGCCC	GCCATAAACT	3240
3241	GCCAGGCATC	AAATTAAGCA	GAAGGCCATC	CTGACGGATG	GCCTTTTGC	GITICIACAA	3300
3301	ACTCTTTTG	TTTATTTTC	TAAATACATT	CAAATATGTA	TCCGCTCATG	AGACAATAAC	3360
3361	CCTGATAAAT	GCTTCAATAA	TATTGAAAAA	GGAAGAGTAT	GAGTATTCAA	CATITCCGIG	3420
3421	TCGCCCTTAT	TCCCTTTTTT	GCGGCATTT	GCCTTCCTGT	TTTTGCTCAC	CCAGAAACGC	3480
3481	TGGTGAAAGT	AAAAGATGCT	GAAGATCAGT	TGGGTGCACG	AGTGGGTTAC	ATCGAACTGG	3540

ATCTCAACAG CGGTA	ΰ	GGTAAGATC	CTTGAGAGTT	TTCGCCCCGA	AGAACGTTTT	CCAATGATGA	3600
GCACTTTTAA AGTTCTGCTA TG	AGTTCTCCTA	TC	rereceese	TATTATCCCG	TGTTGACGCC	GGGCAAGA	3660
AACTCGGTCG CCGCATACAC TAT	CCGCATACAC	TAT	TATTCTCAGA	ATGACTTGGT	TGAGTACTCA	CCAGTCACAG	3720
AAAAGCATCT TACGGATGGC ATG	TACGGATGGC	ATG	ATGACAGTAA	GAGAATTATG	CAGTGCTGCC	ATAACCATGA	3780
GTGATAACAC TGCGGCCAAC TTA	TGCGGCCAAC	TTA	TTACTTCA	CAACGAICGG	AGGACCGAAG	GAGCTAACCG	3840
CTTTTTGCA CAACATGGGG GAT	CAACATGGGG	GAT	GATCATGTAA	CTCGCCTTGA	TCGTTGGGAA	CCGGAGCTGA	3900
ATGAAGCCAT ACCAAACGAC GAG	ACCAAACGAC	GAG	GAGCGTGACA	CCACGATGCC	TGTAGCAATG	GCAACAACGT	3960
TGCGCAAACT ATTAACTGGC GAA	ATTAACTGGC	GAA	GAACTACTTA	CTCTAGCTTC	CCGGCAACAA	TTAATAGACT	402C
GGATGGAGGC GGATAAAGTT GCA	GGATAAAGTT	GCA	GCAGGACCAC	TTCTGCGCTC	GGCCCTTCCG	GCTGGCTGGT	408C
TTATTGCTGA TAAATCTGGA GCC	TAAATCTGGA	OCC	GCCGGTGAGC	GIGGGICICG	CGGTATCATT	GCAGCACTGG	414(
GGCCAGATGG TAAGCCCTCC CGT	TAAGCCCTCC	CGT	CGTATCGTAG	TTATCTACAC	GACGGGGAGT	CAGGCAACTA	4 20C
TGGATGAACG AAATAGACAG ATC	AAATAGACAG	ATC(ATCGCTGAGA	TAGGTGCCTC	ACTGATTAAG	CATTGGTAAC	4 26C
TGTCAGACCA AGTTTACTCA TAT	AGTTTACTCA	TAT	TATATACTTT	AGATTGATTT	AAAACTTCAT	TTTAATTTA	432C
AAAGGATCTA GGTGAAGATC CTT	GGTGAAGATC	CTT	CTTTTTGATA	ATCTCATGAC	CAAAATCCCT	TAACGTGAGT	438C
TITCGITCCA CICAGCGICA GAC	CTGAGCGTCA	GAC	GACCCCGTAG	AAAAGATCAA	AGGATCTTCT	TGAGATCCTT	4440
TITITICIGCG CGTAATCTGC TGC	CGTAATCTGC	TGC	TGCTTGCAAA	CAAAAAAACC	ACCGCTACCA	GCGGTGGTTT	450C
GITTGCCGGA TCAAGAGCTA CCAI	TCAAGAGCTA	CCA	CCAACTCTTT	TTCCGAAGGT	AACTGGCTTC	AGCAGAGCGC	456C
AGATACCAAA TACTGTCCTT CTA	TACTGTCCTT	CTA	CTAGTGTAGC	CGTAGTTAGG	CCACCACTIC	AAGAACTCTG	462C
TAGCACCCC TACATACCTC GCT	TACATACCTC	GCT	GCTCTGCTAA	TCCTGTTACC	AGTECCTECT	GCCAGTGGCG	468C
ATAAGTGGTG TCTTACCGGG TTG	೧೮೮೮	TTG	TTGGACTCAA	GACGATAGIT	ACCGGATAAG	GCGCAGCGGT	474C

5941	CACAATCTTC TCGC	TCGCGCAACG	CGTCAGTGGG	CTGATCATTA	ACTATCCGCT	GGATGACCAG	3009
6001	dardccarre crer	CTGTGGAAGC	TGCCTGCACT	AATGTTCCGG	CGTTATTTCT	TGATGTCTCT	ე909
	GACCAGACAC	CCATCAACAG	TATTATTTC	TCCCATGAAG	ACGGTACGCG	ACTGGGCGTG	6120
	6121 GAGCATCTGG	TCGCATTGGG	TCACCAGCAA	ATCGCGCTGT	TAGCGGGCCC	ATTAAGTTCT	6180
6181	greresees	GICTGCGICT	ggcregcreg	CATAAATATC	rcacregeaa reaaarreag	TCAAATTCAG	624C
6241	CCGATAGCGG AACG	AACGGGAAGG	CGACTGGAGT	GCCATGTCCG	GTTTTCAACA	AACCATGCAA	9300
6301	ATGCTGAATG AGGG	AGGGCATCGT	receacted	Argeregric	ccaaccarca cardecects	GATGGCGCTG	9360
6361	GGCGCAATGC GCGC	GCGCCATTAC	CGAGTCCGGG	CTGCGCGTTG	GTGCGGATAT	CICGGIAGIG	6420
6421	GGATACGACG ATAC	ATACCGAAGA	CAGCTCATGT	TATATCCCGC	CGTCAACCAC	CATCAAACAG	6480
6481		GATITICGCC IGCIGGGCA	AACCAGCGTG	GACCGCTTGC	TGCAACTCTC	TCAGGGCCAG	654 C
6541	GCGGTGAAGG	GCAATCAGCT	GTTGCCCGTC	TCACTGGTGA	TCACTGGTGA AAAGAAAAC	CACCCTGGCG	ე099
6601	CCCAATACGC AAAC	AAACCGCCTC	receeded	TIGGCCGATT	CATTAATGCA	GCTGGCACGA)999
6661	CAGGITICCC GACT	GACTGGAAAG	CGGGCAGTGA	GCGCAACGCA	ATTAATGTGA	GITAGCGCGA	6720
6721	ATTGATCTG						6726